# SPARROW Surface Water Quality Workshop October 29-31, 2002 Reston, Virginia

Section 6. SPARROW Model Calibration

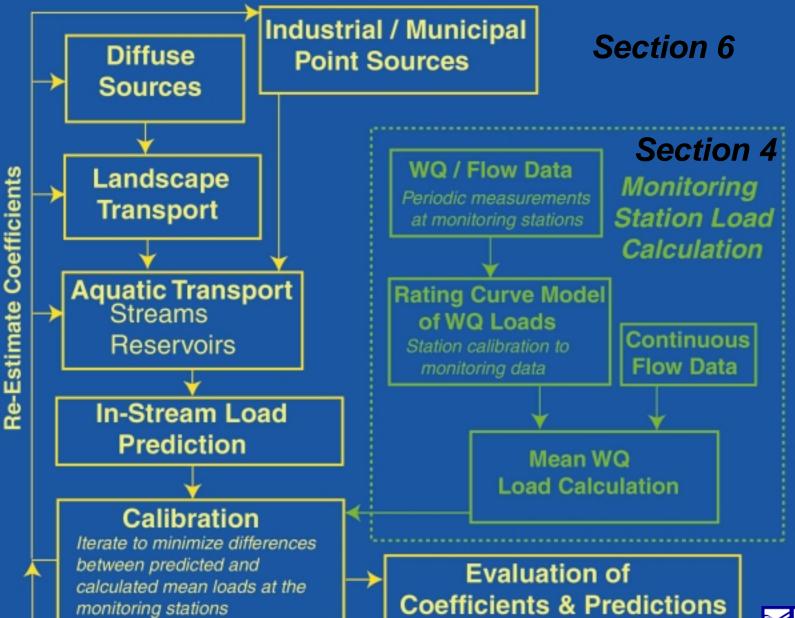


# Section 6. SPARROW Model Calibration Topics Considered

- The SPARROW model equation
- Specification of the model: sources, land-to-water and aquatic transport
- Nonlinear estimation of parameters
- Physical interpretation of parameters
- Evaluating the model error
- Model selection criteria
- SPARROW calibration/prediction software (existing)
   Fecal coliform example
- On-going enhancements to the code

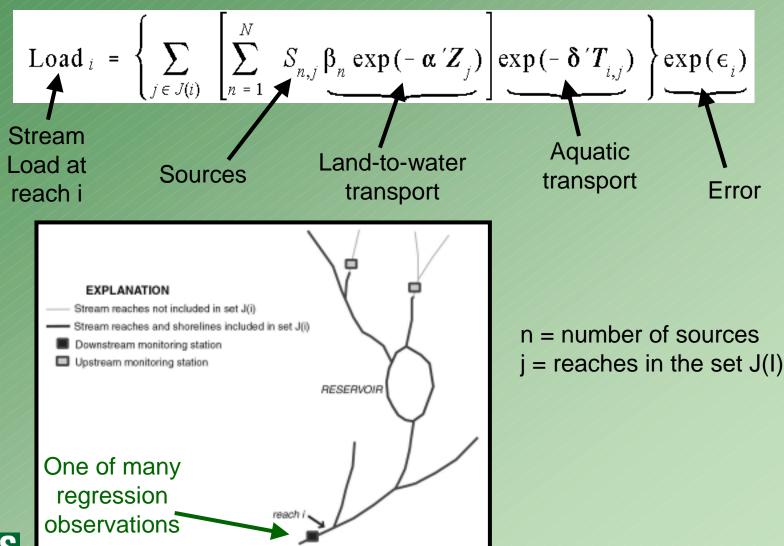


### SPARROW Model Structure





### **SPARROW Model Equation**





### **SPARROW Model Equation**

### Mass-Balance Equation

Load 
$$_{i} = \left\{ \sum_{j \in J(i)} \left[ \sum_{n=1}^{N} S_{n,j} \beta_{n} \exp(-\alpha' Z_{j}) \right] \exp(-\delta' T_{i,j}) \right\} \exp(\epsilon_{i})$$

Stream
Load

Sources

Land-to-water Aquatic transport

Sources

transport

### Model structure is nonlinear:

- Additive sources (preserves mass balance)
- Multiplicative error (account for scale dependency of error)
- Exponential delivery terms



### **Model Equation**

How does the model structure differ from that of a conventional log-linear watershed regression model?

Nonlinear Model: 
$$Y = (X_1B_1 + X_2B_2) e^{\epsilon}$$
  
Log transform:  $ln(Y) = ln(X_1B_1 + X_2B_2) + \epsilon$ 

- Additive sources
- Mass balance
- Multiplicative error

Log-Linear Model: 
$$Y = (X_1^B_1 X_2^B_2) e^{\epsilon}$$
  
Log transform:  $In(Y) = B_1 In(X_1) + B_1 In(X_1) + \epsilon$ 

- Multiplicative sources
- No mass balance
- Multiplicative error



### **SPARROW Model Sources**

Load 
$$_{i} = \left\{ \sum_{j \in J(i)} \left[ \sum_{n=1}^{N} S_{n,j} \beta_{n} \exp(-\alpha' Z_{j}) \right] \exp(-\delta' T_{i,j}) \right\} \exp(\epsilon_{i})$$

Stream
Load

Sources

Land-to-water Aquatic transport

Sources

The stream transport transport

Sources

Sources

Sources

The stream transport

The stream transport

The stream transport

Sources

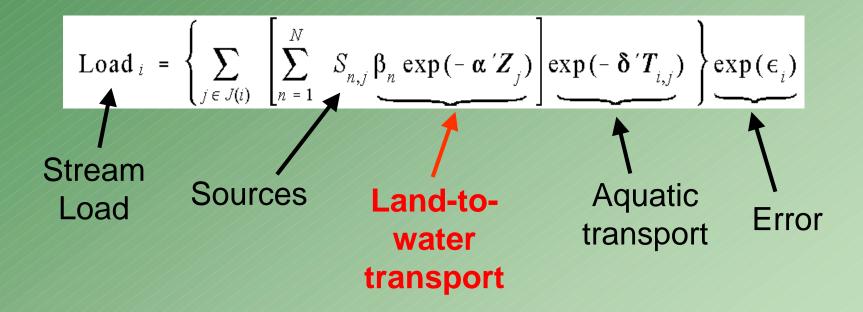


### **SPARROW Model Sources**

- Selection of sources determined by:
  - Research literature
  - Your expertise knowledge of the watershed
  - Data availability
- Diffuse Sources:
  - Extensive (land-use)
  - Intensive (mass based)
  - Mixed model (extensive and intensive)
- Point sources:
  - Contaminant mass (expect coefficient of 1.0 if response variable in same units and model properly specified)
  - Surrogates: sewered population; BOD; Flow
- Geographic dummy variables (e.g., unspecified sources)



### **SPARROW Landscape Transport**





### **SPARROW Model Landscape Transport**

### Selection of landscape variables:

- Variables should relate closely to landscape processes (e.g., runoff and drainage area integrate terrestrial and aquatic processing and water transport—not recommended)
- Water balance inputs and landscape-related loss components (precipitation; evapotranspiration)
- Soil properties (e.g., organic content, permeability, moisture content)
- Water flow paths (e.g., TOPMODEL overland flow; DEM overland routing)
- Management activities (e.g., tile drainage, conservation tillage practices; BMPs—stream riparian properties)
- Land use (e.g., wetlands—is it a source or sink?; measures of impervious surface or urbanization may confound source estimation)



### **SPARROW Model Landscape Transport**

### Landscape decay functions:

- Exponential function with imbedded negative sign: constrains values between zero and one:
  - -- All coefficients reported with positive sign
  - -- Negatively related variables:

$$e^{(-\beta_1 X_1 - \beta_2 X_2)}$$
 (e.g., soil permeability)

--Positively-related variables enter as reciprocal:

$$e^{(-\beta_1^{1/X}_1 - \beta_2^{1/X}_2)}$$
 (e.g., drainage density)

 Exponential function with log transformed variables – unconstrained:

$$e^{(\beta_1 \log(X_1) + \beta_2 \log(X_2))}$$

-- Coefficients reported with actual sign



### **SPARROW Model Landscape Transport**

### Landscape-source variable interactions:

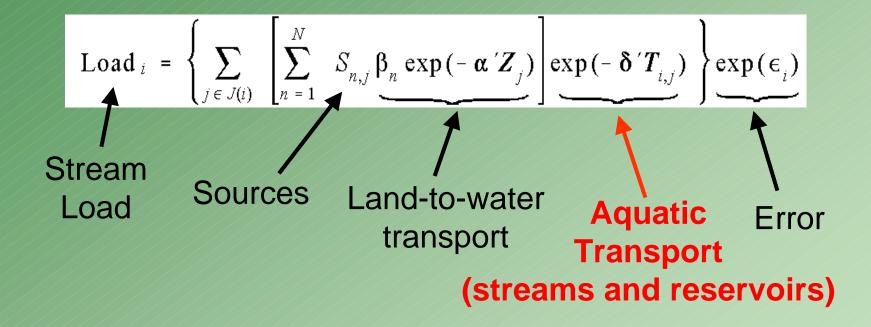
 Values of diffuse-source coefficients not independent of delivery variables (interaction not separable in model)

Load<sub>i</sub> = 
$$\left\{ \sum_{j \in J(i)} \left[ \sum_{n=1}^{N} S_{n,j} \beta_{n} \exp(-\alpha' Z_{j}) \right] \exp(-\delta' T_{i,j}) \right\} \exp(\epsilon_{i})$$

- Standardize delivery variables (deviations from mean) to create more interpretable source coefficients expressed in relation to mean delivery—yields reflecting delivery to streams best metric to reflect geographic variations
- Commonly assume that diffuse sources subject to same landscape and aquatic decay (however, code can accommodate separate source-delivery interactions)



### **SPARROW Aquatic Transport**





### **Stream Transport**

### **Chemical Reaction Kinetics**

-The reaction rate is proportional to the concentration of the

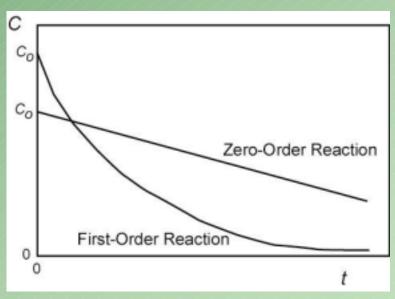
reactants (law of mass action):

$$dc / dt = -kc^n$$

 $k = loss rate (T^{-1})$ 

c = concentration of a single reactant

n = reaction order



-Integrated equation for the <u>first-order model</u> (exponential depletion):

$$c = c_0 e^{-kt}$$



$$Flux = Flux_0 e^{-kt}$$

Loss rate (per unit water travel time) integrates multiple processes:

$$k = k_D + k_U + k_S$$

where,

 $k_D$  = denitrification rate (N only) ~ f[benthic areal rate(+), depth(-), concentration(+/-), temperature(+), organic matter(+), flow(-)]

 $k_U$  = biological uptake rate ~ f[algal density (+), light(+), depth(-), concentration(+), temperature(+)]

 $k_S$  = settling loss rate ~ f[particle settling velocity (+), depth(-), particulate bound fraction (+)] (particulate burial?)

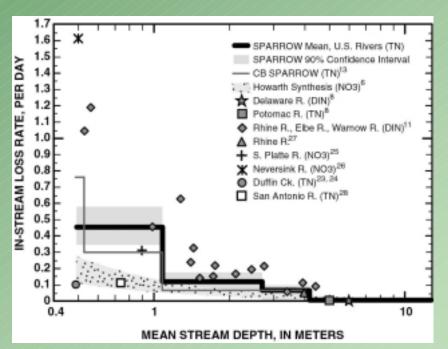
t = water time of travel

Related Questions: time scale of steady state processes (multiple species model) and spatial scaling factors

Depth (streamflow) theoretically important spatial scaling factor affecting nutrient loss rate:

$$Flux = Flux_0 e^{-k_n t}$$

where, n = flow class defined according to mean streamflow

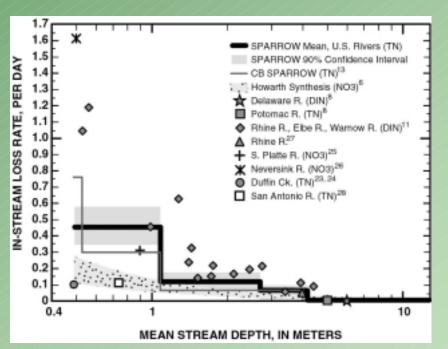


SPARROW
Estimates of Total
Nitrogen Loss vs.
Channel Depth



### Approach for defining flow classes:

- Use a few integer factor and order of magnitude separations
- -Start simple with 2 to 3 flow classes
- -Evaluate final selections with continuous flow functions



SPARROW
Estimates of Total
Nitrogen Loss vs.
Channel Depth



Per unit time (day-1) rate: Flux = Flux<sub>0</sub>  $e^{-k}$ <sup>t</sup>

Per unit channel length (km<sup>-1</sup>) rate: Flux = Flux<sub>0</sub>  $e^{-k'}$ <sub>n</sub> Flux = Flux<sub>0</sub>  $e^{-k'}$ <sub>n</sub>

### Approaches to obtaining day-1 rate:

- Post-conversion:  $km^{-1}$  to day<sup>-1</sup> rate using available estimates of mean stream velocity ( $k = k'^*$  V) for comparison with literature values
- Pre-conversion: Stream morphological / hydrology studies (e.g., Leopold and Maddock, 1953; Jobson, 1996; Jowett, 1998) can be used to estimate time-of-travel for individual reaches based on streamflow

### Obtaining estimates of channel depth:

e.g., Leopold and Maddock, 1953; Depth = 0.2612 Q 0.3966



### **Nutrient Transport in Reservoirs/Lakes**

### **Empirical mass-balance models:**

- Vollenweider, 1969
  - Phosphorus: Reckhow and Chapra, 1983
  - Nitrogen: Kelly et al. 1987; Molot & Dillon, 1993
- Steady state, well-mixed conditions
- Retention ~ f(depth, residence time, volume, areal water load, and apparent settling velocity—mass transfer coefficient)



### **Lake Retention of Nutrients**

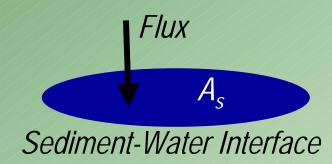
Depth-Independent vs. Depth-Dependent Approaches

Constant Settling Velocity: Settling =  $v A_s c$ 

v = apparent settling velocity (units: L T<sup>-1</sup>)

 $A_s$  = lake surface area

c = concentration



First-order reaction (depth-dependent settling velocity):

Reaction =  $k_s$  V c

 $k_s = v / \text{depth}$ ; first-order settling rate (units=T<sup>-1</sup>)

V = volume (units=L<sup>3</sup>=depth\*surface area)

### Which is better?

- Equivalent mathematically
- Former more specific to process of flux across sediment-water interface
- Little empirical evidence in lake literature of detectable differences
- Availability of data may determine (volume vs. surface area)



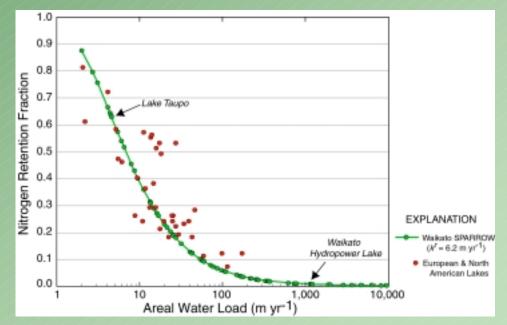
### **Nutrient Transport in Reservoirs/Lakes**

### **SPARROW** estimated loss:

- Settling velocity coef. (constant in all water bodies)
- Areal water load (ratio of mean outflow to surface area)



SPARROW Estimates of Total Nitrogen Loss (75 Reservoirs of the Waikato River Basin, New Zealand)





# Lake Retention of Nutrients Current SPARROW Equation

From Mass-Balance Expression (Depth Independent) (e.g.,Reckhow and Chapra, 1983)

$$R = v / (v + q_s)$$

R = retention coefficient v = apparent settling velocity (L T<sup>-1</sup>)  $q_s$  = areal water load (outflow/surface area; L T<sup>-1</sup>)



# Lake Retention of Nutrients Previous SPARROW Equation

**Empirical Approximation to Mass-Balance Expression** 

(Alexander et al. Wat. Resour. Res., in press)

$$R = 1 - exp(-v/q_s)$$

R = retention coefficient v = apparent settling velocity (L T<sup>-1</sup>)  $q_s =$  areal water load (L T<sup>-1</sup>)



# Lake Retention of Nutrients Original SPARROW Equation

(Smith et al. Wat. Resour. Res., 1997)

$$R = 1 - exp(-kT)$$

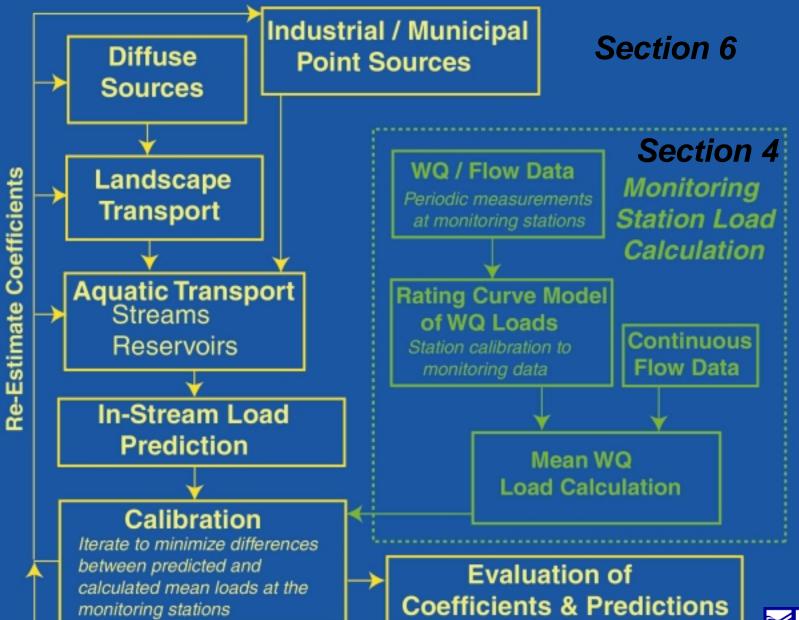
R = retention coefficient

k = 1<sup>st</sup>-order decay rate

T = channel water travel time in reservoir reach



### SPARROW Model Structure





## SPARROW Model Equation Nonlinear Estimation of Parameters

Load<sub>i</sub> = 
$$\left\{ \sum_{j \in J(i)} \left[ \sum_{n=1}^{N} S_{n,j} \beta_{n} \exp(-\alpha' Z_{j}) \right] \exp(-\delta' T_{i,j}) \right\} \exp(\epsilon_{i})$$
Stream
Load
Source
Land-to-water
parameters
parameters

Error

### Model structure is nonlinear:

- Additive sources (preserves mass balance)
- Multiplicative error (account for scale dependency of error)
- Delivery terms



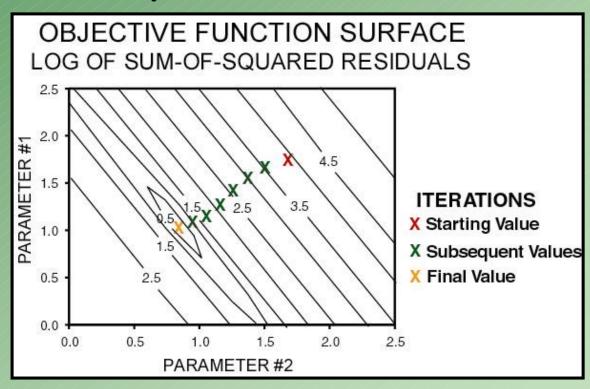
Nonlinear Regression can be viewed as an extension of linear regression analysis.

Uses <u>Gauss-Newton optimization method</u> (Levenberg-Marquardt conditioning parameters)—an iterative form of standard linear regression

- User selects starting values for the parameters.
- Method iteratively applies linear approximations of the nonlinear model in the vicinity of the initial and subsequent parameter values until the model <u>converges</u>.
- Model <u>converges</u> when changes in the parameter estimates are less than a preset threshold. The parameters minimize the <u>objective</u> <u>function</u>.
- Objective function is a measure of the fit between predicted and observed values (e.g., sum-of-squared residuals)



- The optimization routine uses a linear approximation of the nonlinear objective function.
- Parameter values are changed iteratively to locate the minimum value of the objective function.





### How do you pick starting values?

- Literature values
- Other SPARROW models or local models
- Guestimate remaining values if lack information

### Model convergence:

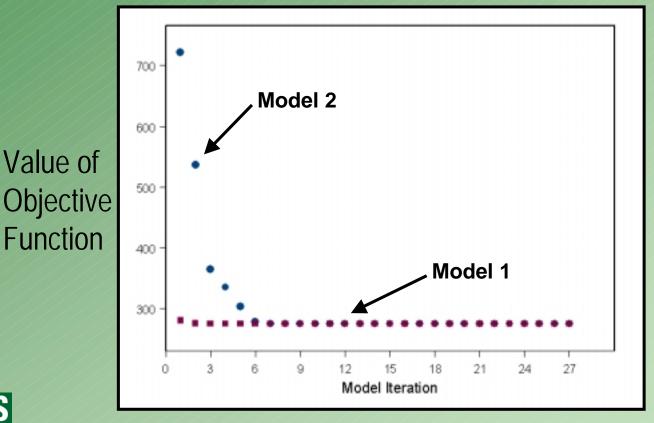
- If model well conditioned, only exceedingly large differences between starting and final values will be problematic
- Should test the convergence of final model for stability (selection of global rather than local minima) by changing starting values by a large amount
- SPARROW convergence problems often related to data errors; problems can occur when attempt to estimate too many parameters for certain functions (e.g., reservoir decay)



### Model Iterations of Fecal Coliform Model

Model 1: Starting values near final estimates (33 iterations)

Model 2: Starting values changed by 2 orders magnitude (aquatic decays by 1 order magnitude) – (27 iterations)



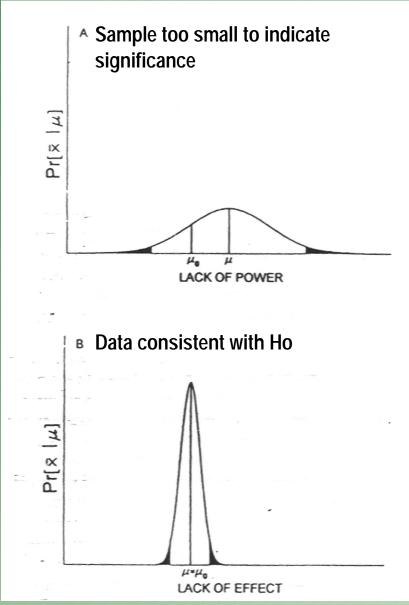
Final parameter estimates identical for two models



### Check if converged model reasonable:

- Do the parameter estimates have the correct sign?
- Are the parameters and standard errors statistically significant?
  - t-statistics: t = b<sub>j</sub> / se(b<sub>j</sub>)
  - tests of parameter significance:
    - t values approximate (only asymptotically valid)
    - strict adherence to α level not recommended
    - insignificant parameters: "lack of effect" vs. "lack of power"





# Lack of Power vs. Lack of Fit

Possible Interpretations of a Test That Fails to Reject Ho (e.g., p>0.05)

From: Johnson, 1999, J Wildl. Manage., v. 63



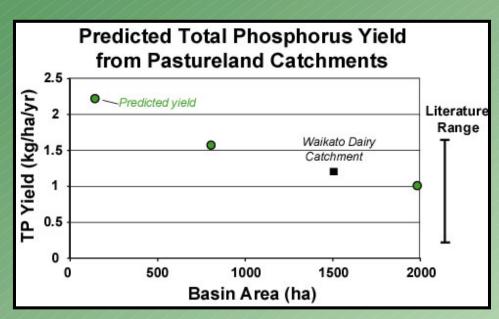
### Check if converged model reasonable:

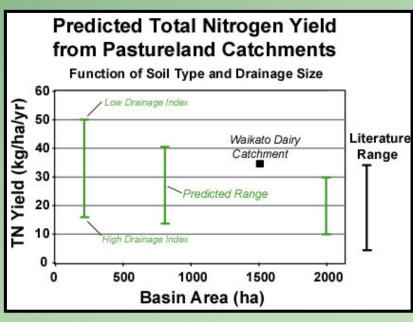
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  - tests of parameter significance:
    - t values approximate (only asymptotically valid)
    - strict adherence to α level not recommended
    - insignificant parameters: "lack of effect" vs. "lack of power"
- Do the parameters have physical significance?

Literature comparisons: catchment yields by land use, per capita waste loads, point-source coef., in-stream decay, reservoir settling rates



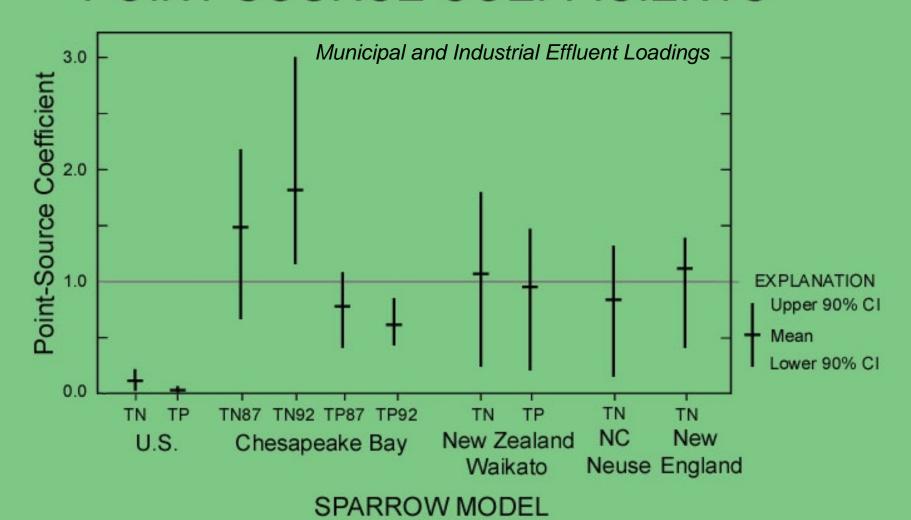
# Verification of Estimated Diffuse Source Coefficients New Zealand SPARROW







## NATIONAL & REGIONAL SPARROW POINT-SOURCE COEFFICIENTS

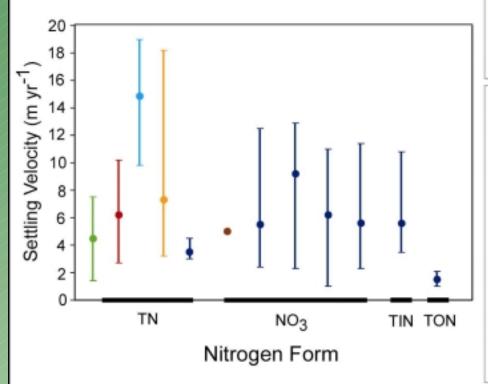




### SPARROW Estimates of Nitrogen Loss in Reservoirs

Mean & Range for Nitrogen Settling Velocity

Rates for Lakes & Reservoirs\*



#### SPARROW Models

U.S. (1992 preliminary)

#### N.Z. Waikato

Alexander et al. (in press)

N.C. Neuse Coastal

#### LITERATURE DATA

#### Danish Lakes

Windolf et al. (1996)

#### S. Ontario Lakes

Molot & Dillon (1993)

Kelly et al. (1987)

Kelly et al. (1990)

Dillon & Molot (1990)

#### Lake Superior

Bennett (1986)

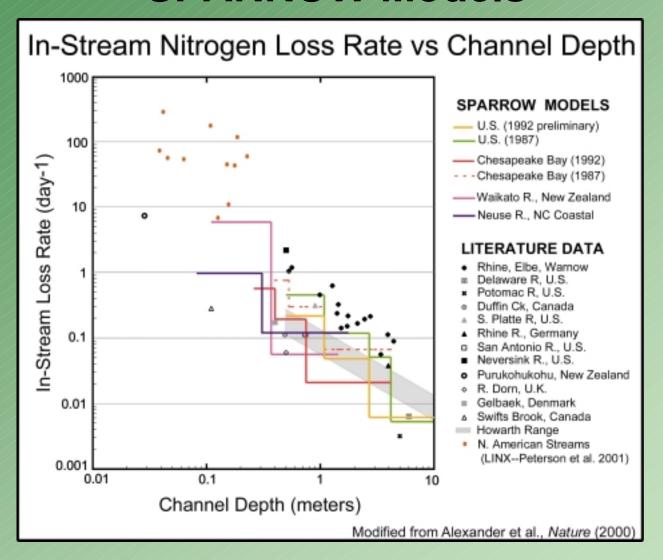
#### Common Attributes:

High N inputs Mod. to high N:P ratios Denitrification dominated

\* Settling rates > 25 m yr<sup>-1</sup> - algal dominated; low N:P ratios (Kelly et al. 1990)



# National and Regional SPARROW Models





### **Nonlinear Parameter Estimation**

#### Check if converged model reasonable:

- Do the parameter estimates have the correct sign?
- Are the parameters and standard errors statistically significant?
  - t-statistics: t = b<sub>j</sub> / se(b<sub>j</sub>)
  - tests of parameter significance:
    - t values approximate (only asymptotically valid)
    - strict adherence to α level not recommended
    - insignificant parameters: "lack of effect" vs. "lack of power"
- Do the parameters have physical significance?
   Literature comparisons: catchment yields by land use, per capita waste loads, point-source coef., in-stream decay, reservoir settling rates
- Are the parameters correlated?



## Parameter Correlation: Multicollinearity

#### What's the problem?

- Coefficient signs unreasonable
- Two variables describing the same process have different signs and insignificant coefficients

#### Metrics to detect it:

- Parameter correlation matrix very high correlations (>0.95)
- Variance Inflation Factors (VIFs) measure of correlation among all explanatory variables (>10 a problem)

#### How do you fix it?

- Center data fix for polynomial variables
- Simplify model
- Remove parameter
- Combine data (equating two parameters)



### **Nonlinear Parameter Estimation**

#### Model convergence can also be assisted by:

- Log transformations of data and parameters
   May be needed to satisfy residual assumptions (linearity, constant variance, normally distributed)
- <u>Scaling of data</u> better conditioning of the derivative matrix if reduce large differences in coefficient values:

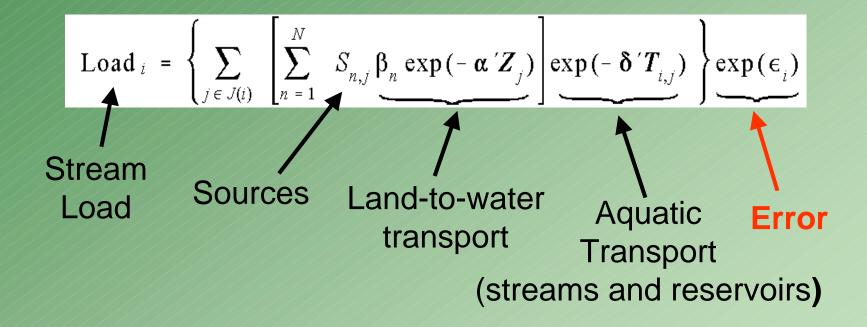
$$Y = b0 \exp(-b1 X)$$

If expect b0=100 and b1=0.001, force coefficients to be approx 1.0:

$$Y = 100 b0 exp(-b1 X/1000)$$

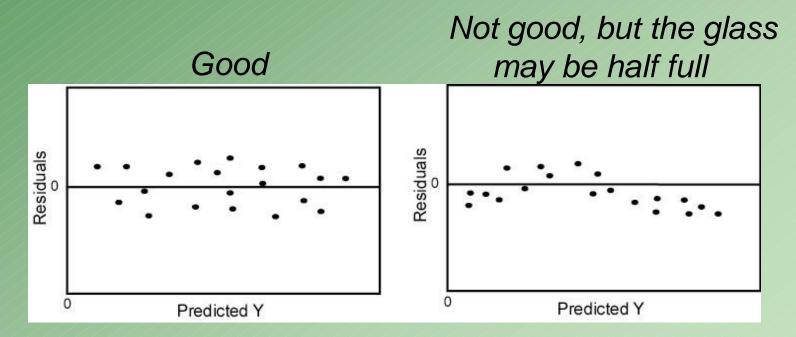


#### **SPARROW Error Term**





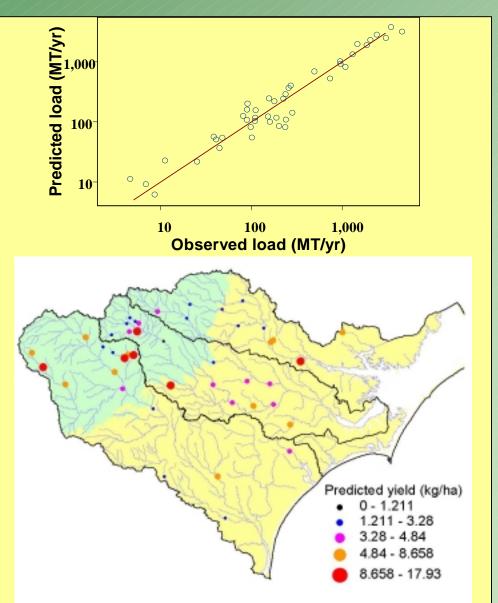
 Residuals should vary randomly – no evidence of systematic patterns that may indicate correlation or explanatory variables missing from the model



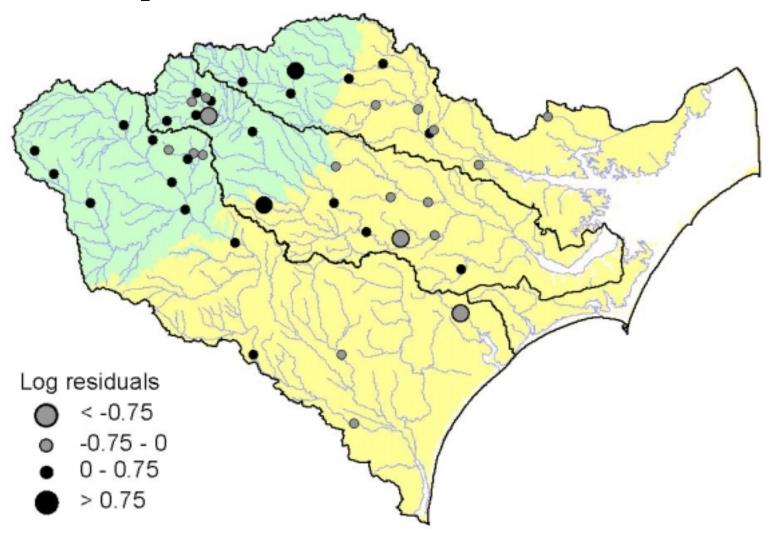


## NC SPARROW land cover-based regression model

R2	0.93	
MSE	0.22	
TN sources		
Point sources (MT/1992)	Parameter 0.85	p-value <b>0.006</b>
Agr. Land area (MT/km²)	<b>5.9</b>	0.09
Non-agr land area (MT/km²)	1.79	0.08
Land delivery variable		
Soil hydrologic group	4.13	0.001
Aquatic loss		
Small stream (km <sup>-1</sup> )	0.08	0.02
Large stream (km <sup>-1</sup> )	0.002	0.35
Reservoir (m/yr)	16.4	0.008



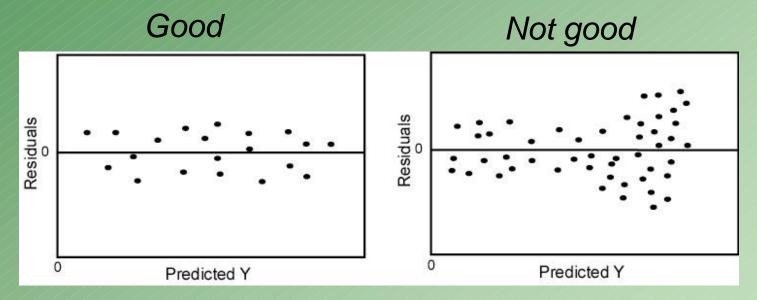
#### **Spatial distribution of model residuals**





Assumptions affecting accuracy of parameter standard errors:

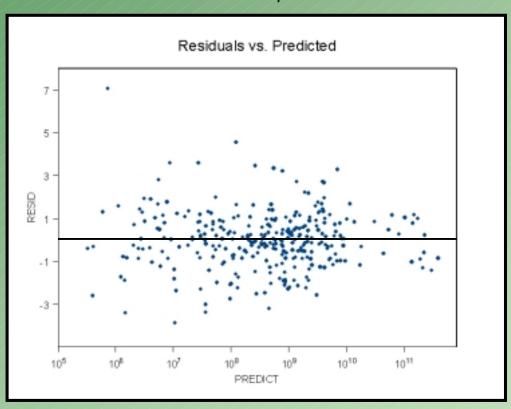
Variance of the residuals should be constant (homoscedastic)



<u>Solution</u>: Weighting of residuals for measurement error (e.g., load estimation error) and model variance—assumes error-variance relations can be defined.



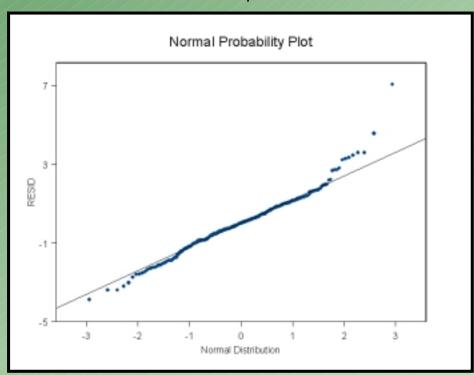
Fecal Coliform Residual Plot 305 NASQAN sites, records 1978-92





Residuals are normally distributed

Fecal Coliform Residual Probability Plot 305 NASQAN sites, records 1978-92



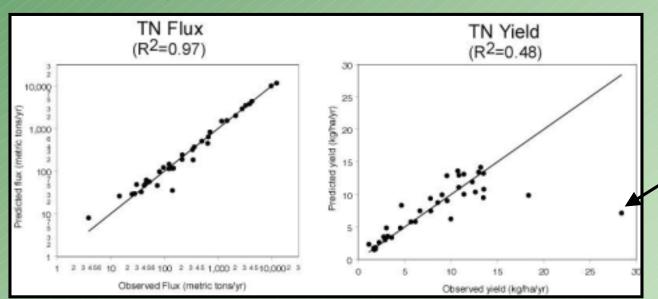


#### **Outliers**

"I like SPARROW's ability to identify inaccuracies in monitoring data...the model is right and the data are wrong."

-Graham McBride, NIWA, NZ

Caused by data errors or model mis-specification



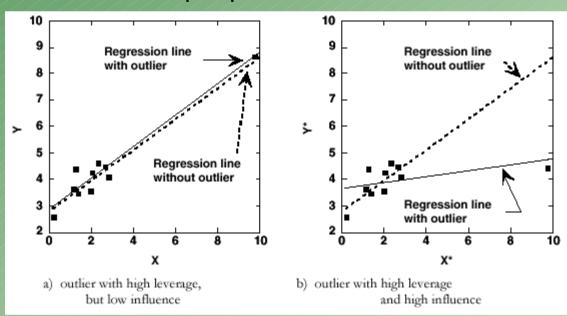
#### NZ Waikato TN Model

Horticulture prominent in the watershed— source unspecified in the model



#### **How Detect Outliers in Multi-Dimensional Space?**

- <u>Leverage</u> (hat matrix)—measure of outlier in at least one of the explanatory variables
  - --High leverage for value > 3p / n, where p=#parameters; n=# observations



- Influence statistics

   (outliers in the response variable):
- --Standardized residuals (standard deviation units)
  - --Cook's D



### Significance of model parameters – nested models

- Individual parameters t statistics (and associated p values)
- Multiple parameters F test

The test statistic is 
$$F = \frac{(SSE_s - SSE_c) / (df_s - df_c)}{(SSE_c / df_c)} \qquad \text{where } (df_s - df_c) = m-k.$$



### Significance of model parameters – nested models

- Individual parameters t statistics (and associated p values)
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SSy

SSR (signal)

SSE (noise)

```
Total sum of squares = Treatment sum of squares (overall variation) = (group means – overall mean) + (variation within groups)
\sum_{j=1}^{k} \sum_{i=1}^{n_j} (y_{ij} - \overline{y})^2 = \sum_{j=1}^{k} n_j (\overline{y}_j - \overline{y})^2 + \sum_{j=1}^{k} \sum_{i=1}^{n_j} (y_{ij} - \overline{y}_j)^2
```



### Overall model fit

- <u>Lowest MSE</u> (Mean Square Error)—declines with increasing # parameters: MSE = SSE / (n-p)
- <u>Highest R-squared</u>—increases with increasing # parameters

$$R^2 = 1 - (SSE / SS_y)$$

where,  $SS_y$  = total sum of squares of the regression equation SSE = sum of square error

 Highest Adjusted R-squared (adjusted for # parameters; relatively insensitive when n>>p):

$$R^2 = 1 - [(n-1) SSE] / [(n-p) SS_y]$$



### Overall model fit

- Lowest Mallow's Cp (Cp adjusted for # parameters)
- Lowest PRESS statistic
- Your professional judgement

$$Cp = p + \frac{(n-p) \cdot (s_p^2 - \hat{s}^2)}{\hat{s}^2}$$

where Sp = MSE of the p parameter model s2 = best estimate of the "true" error (lowest MSE of all models)



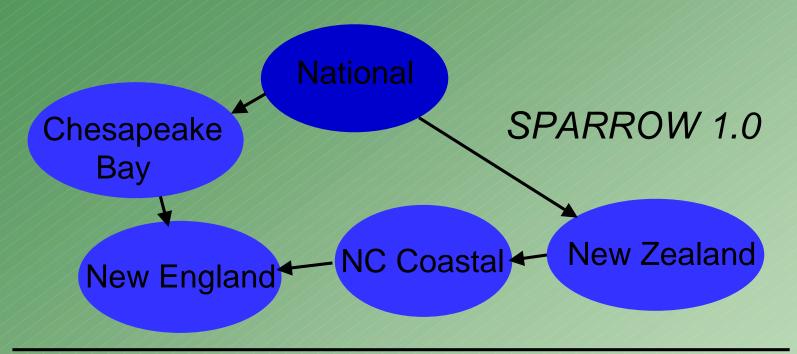
# SPARROW Model Calibration Summary

#### Evaluation of model structure and fit

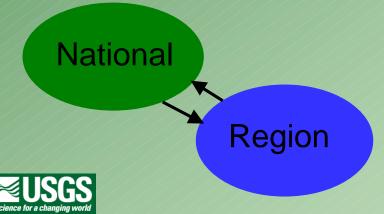
- 1. Check parameter estimates for statistical and physical significance, correct sign, correlation, and stability
- 2. Check residual plots for outliers, systematic patterns, homoscedasticity, normality, and inspect mapped residuals
- 3. Check overall model fit (adjusted R-squared, MSE, Mallow's Cp)



#### **Evolution of SPARROW Calibration Software**



#### SPARROW 2.0



#### SAS calibration / prediction software

- New revisions completed and on-going to national code
- Document and support single source of software maintained by the national SPARROW group

GIS methods and software –variety of approaches—will likely continue

#### Preprocessing steps:

- SPARROW explanatory variables identified by reach using GIS
- Network navigation parameters (hydrologic order of reaches, from- and to-nodes, diversion fraction)
- Assembly of SAS data set (DATA1)
  - Compiles reach-level data from GIS processing in single dataset
- SAS model procedure for calibration and prediction
  - Initial parameter setup (modify variable lists)
  - Data block
    - Final calculations
    - Data screening (e.g., stations)
  - IML procedures (landscape and aquatic decay equations)
  - Output parameters and predictions



# Fecal Coliform Bacteria Example Model Building Exercise

#### Possible sources

Human and animal sources—wastewater, urban runoff, and septic systems, livestock populations (confined feeding operations, unconfined), background for other animals (geese, birds, etc.)

#### Loss rate for total coliform bacteria:

$$k_T = k_B + k_R + k_S$$

where,

 $k_B$  = base mortality rate (fresh waters) = 0.8 \* 1.07 T-20

k<sub>R</sub> = solar radiation effect ~ f[light energy (+), depth(-), particulate matter (-)]

k<sub>S</sub> = settling loss rate ~ f[particle settling velocity (+), depth(-),
fraction of attached bacteria(+)]



## **SPARROW Fecal Coliform Models**

## **Intensive-source model:**

Livestock wastes (confined and unconfined), Sewered population, Urban land, Other lands

## Land-use (extensive-source) model:

Agricultural lands, Sewered population, Urban land, Other lands



1. Define response (dependent) and source variables

```
/* Specify all the variable lists. */

/* Dependent variable */
%let depvar = load;

/* Source variables */
%let srcvar = SEWERPOP RESLAND CONF UNCONF URBAN;

/* Source variable coefficients */
%let bsrcvar = BPOINT BRESLAND BCONF BUNCONF BURBAN;
```

SEWERPOP = sewered population

CONF = confined feeding wastes (kg N)

UNCONF = unconfined feeding wastes (kg N)

URBAN = urban land area (km2)

RESLAND = other land (forest, barren, wetlands, shrub)

area (km2)



2. Define landscape delivery variables (constrained exponential function)

```
/* Delivery variables */
%let dlwar = aperm aidrainden ;

/* Delivery variable coefficients */
%let bdlwar = bperm bdrainden ;
```

APERM = Soil permeability
(mean adjusted)

AIDRAINDEN = Drainage density
(reciprocal mean adjusted)

Adjusted delivery variables (user prompt in future version)

```
/* adjust land-to-water delivery factors by mean */
PROC MEANS DATA=indata;
VAR perm idrainden;
OUTPUT OUT=mean_ltw MEAN= xperm xdrainden;
RUN;

data indata; if _n_ = 1 then set mean_ltw; set indata;
aperm = perm - xperm;
aidrainden = idrainden - xdrainden;
run;
```

PERM = Soil permeability IDRAINDEN = Drainage density (reciprocal)



#### 3. Define aquatic decay variables

```
/* Decay variables */
%let decvar = rchtot1 rchtot2 rchtot3;

/* Decay variable coefficients */
%let bdecvar = brchtot1 brchtot2 brchtot3;

/* Reservoir variables */
%let resvar = iresload;

/* Reservoir variable coefficients */
%let bresvar = bresload;
```

```
RCHTOT1 = Reach TOT (days; flow <100cfs)
RCHTOT2 = Reach TOT (days; flow 100 to 500 cfs)
RCHTOT3 = Reach TOT (days; flow >500cfs)

IRESLOAD = Areal hydraulic load (reciprocal; yr/m)
for reservoir outlets
```

#### Create flow interval variables in data section

```
RCHTOT1 = (meanq <= 100) * (rchtype = 0) * RCHTOT;
RCHTOT2 = (100 < meanq <= 500) * (rchtype = 0) * RCHTOT;
RCHTOT3 = (meanq > 500) * (rchtype = 0) * RCHTOT;

if RHLOAD ^= . and rchtype = 2 then iresload = RHLOAD;
else iresload = 0;
```

```
MEANQ = mean streamflow (cfs)

RCHTYPE = reach type code

0=river reach

1=reservoir reach

2=reservoir outlet
```



4. Define delivery variable design matrix

#### **DLVDSGN** Code

0 = delivery not apply to this source

1 = delivery applies to this source

/\* Specify the delivery design matrix: each row is a different source (in the same order as they are listed in the srcvar statement); each column is a different delivery variable (in the same order as they are listed in the dlvvar statement). An element is either a 0 or 1. Element r,c is a 1 if source r uses delivery variable c. Otherwise, element r,c is 0. A space separates columns and a comma separates rows. \*/

%let dlvdsgn = 0 0, 1 1, 1 1, 1 1, 1 1;

Displayed sequence for DLVDSGN: 0 0 = sewered population for permeability and drainage density

1 1 = residual land area for permeability and drainage density ...etc.



#### 5. Select estimation and/or prediction execution mode

```
%let if_estimate = yes;
%let if_predict = yes; * Specify if predictions are to be made.;
%let if_adjust = no ; * Specify if the load predictions (decayed) are to be adjusted for actual loads at monitoring stations;
```

6. Define data set, hydrologic sorting variable, and parameter starting values



7. Define reach and station IDs and network navigation parameters (set once—no need to change)

```
/* Assign a list of column reference vectors */
%let makecol =
   jwaterid = %col(datalst,waterid) %str(;)
   jstatpk = %col(datalst,statpk) %str(;)
   jfnode = %col(datalst,fnode) %str(;)
   jtnode = %col(datalst,tnode) %str(;)
   jfrac = %col(datalst,frac) %str(;)
   jaiftran = %col(datalst,aiftran) %str(;)
```

WATERID = watershed / reach ID

STATPK = station ID

FNODE = reach upstream node

TNODE = reach downstream node

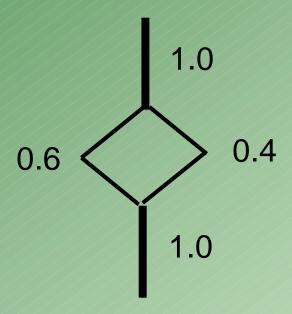
FRAC = reach diversion fraction

AIFTRAN = transport flag (1=transport reach)

```
/* Make list of variables to be read from the SAS indata data set and loaded
  into a matrix. Detect and remove duplicates. */
%let addlist = &depvar &srcvar &dlvvar &decvar &resvar &othvar;
%let datalst = waterid statpk tnode fnode frac aiftran;
```



FRAC - Fraction diversion value





7. Define reach and station IDs and network navigation parameters (set once—no need to change or transparent if use same names)

```
/* Assign a list of column reference vectors */
%let makecol =
   jwaterid = %col(datalst,waterid) %str(;)
   jstatpk = %col(datalst,statpk) %str(;)
   jfnode = %col(datalst,fnode) %str(;)
   jtnode = %col(datalst,tnode) %str(;)
   jfrac = %col(datalst,frac) %str(;)
   jaiftran = %col(datalst,aiftran) %str(;)
```

WATERID = watershed / reach ID STATPK = station ID FNODE = reach upstream node TNODE = reach downstream node FRAC = reach diversion fraction AIFTRAN = transport flag (1=transport reach)

```
/* Make list of variables to be read from the SAS indata data set and loaded
  into a matrix. Detect and remove duplicates. */
%let addlist = &depvar &srcvar &dlvvar &decvar &resvar &othvar;
%let datalst = waterid statpk tnode fnode frac aiftran;
```

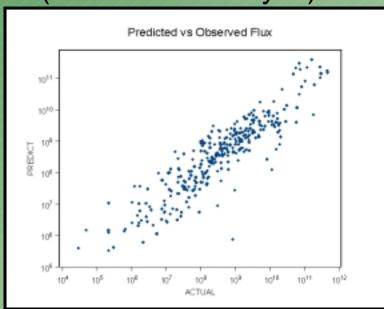


#### INTENSIVE MODEL RESULTS: Fecal Coliform

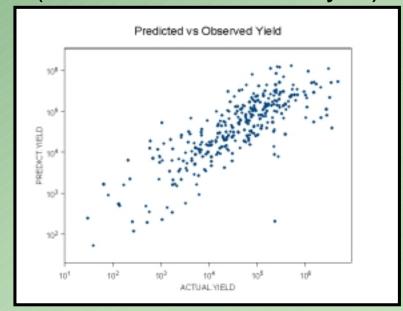
305 NASQAN sites, records 1978-92

Sources reflect mean adjusted landscape variables

Coliform Flux (10<sup>-2</sup> Bcolonies yr<sup>-1</sup>)



Coliform Yield (10<sup>-2</sup> Bcolonies km<sup>-2</sup> yr<sup>-1</sup>)



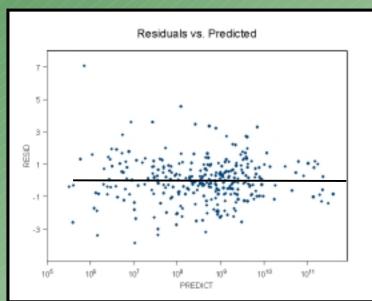


#### INTENSIVE MODEL RESULTS: Fecal Coliform

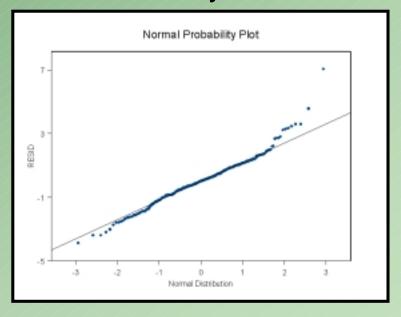
305 NASQAN sites, records 1978-92

Sources reflect mean adjusted landscape variables

#### Residual Plot



#### **Probability Plot**





#### INTENSIVE MODEL RESULTS: Fecal Coliform

305 NASQAN sites, records 1978-92

Sources reflect mean adjusted landscape variables

Non-linear Least Squares Results							
N Obs DF M 305	odel DF E		SSE 3903 1.867		MSE R-Squ 7443 0.8071	_	R-Sq )0574
	Parameter	Estimate	Std Err	t Value	Pr >  t	Coeffic	eient Units
Sources							
Sewered Population	BPOINT	8919.403	2958.4093	3.0149321	0.0027946	8,919/10	00 = 89.2 Bcol/person/yr
Residual Land	BRESLAND	2045.9055	4741.886	0.4314539	0.6664545	20 Bcol/	km2/yr (0.20 Bcol/ha/yr)
Confined wastes	BCONF	324.97938	116.48886	2.7897894	0.0056188	3.25 Bco	l/kg N/yr
Unconfined wastes	BUNCONF	177.07559	73.883056	2.3967009	0.0171679	1.77 Bco	l/kg N/yr
Urban land	BURBAN	3444100.3	2068444.4	1.6650678	0.0969645		Scol/km2/yr Scol/ha/yr)
Landscape loss							
Soil permeability	BPERM	0.334435	0.0841099	3.976165	0.0000882	h/cm	(inverse relation)
Drainage density	BDRAINDEN	0.053385	0.0393925	1.3552088	0.1763913	per km	(positive relation)
Aquatic loss							
Stream decay(<100cfs)	BRCHTOT1	0.6485695	0.2210388	2.9341881	0.0036077	per day	
Stream decay(100-500cfs)	BRCHTOT2	0.4942857	0.1846064	2.6775113	0.007834	per day	
Stream decay(>500cfs)	BRCHTOT3	0.1270183	0.0568133	2.2357144	0.026121	per day	
Reservoir decay	BRESLOAD	78.995344	22.727153	3.4758134	0.0005861	m/day	



#### EXTENSIVE (LAND-USE) MODEL RESULTS: Fecal Coliform

305 NASQAN sites, records 1978-92

Sources reflect mean adjusted landscape variables

#### Non-linear Least Squares Results

```
N Obs DF Model DF Error
                                            SSE
                                                      MSE Root MSE R-Square Adj R-Sq
               305
                                   295 573.11607 1.9427663 1.3938315 0.7987313 0.7925909
                         Parameter Estimate
                                              Std Err
                                                        t Value Pr > |t|
                                                                              Coefficient Units
Sources
Sewered Population
                        BPOINT
                                   9345.4095 3061.8093 3.0522507 0.0024781
                                                                            9.345/100 = 93.5 Bcol/person/vr
Cultivated Land
                        BAGRIC
                                   603793.17 185355.63 3.257485 0.0012552
                                                                            6,038 Bcol/km2/yr (60.4 Bcol/ha/yr)
Residual Land
                        BRESLAND
                                                                            68.7 Bcol/km2/yr (0.69 Bcol/ha/yr)
                                   6870.043 3741.5327 1.8361574 0.0673406
                                                                            42,514 Bcol/km2/yr (425.1 Bcol/ha/yr)
Urban land
                        BURBAN
                                   4251364.3 2457871.4 1.7296936 0.084731
Landscape loss
Soil permeability
                        BPERM
                                   0.2356181 0.0788437 2.9884218
                                                                   0.00304
                                                                           h/cm
                                                                                     (inverse relation)
Drainage density
                        BDRAINDEN 0.1459664 0.0786509 1.8558782 0.0644681 per km
                                                                                     (positive relation)
Aguatic loss
Stream decay(<100cfs)
                        BRCHTOT1
                                   0.6405845 0.2213565 2.8939041 0.0040886
                                                                            per day
Stream decay(100-500cfs) BRCHT0T2
                                   0.5339951 0.1996834 2.6742096 0.0079082
                                                                            per day
                        BRCHTOT3
Stream decay(>500cfs)
                                    0.126373 0.0601968 2.099331 0.0366373
                                                                            per dav
Reservoir decay
                        BRESLOAD
                                   70.624909 21.608323 3.2684123 0.0012094
                                                                            m/dav
```

Human fecal coliform intestinal bacteria = 730 Bcol/person/yr Cultivated land = NLCD pasture + row crops + fallow land + orchards



### **Land-Use Model**

## **Intensive-Use Model**

+/- 137%

Sources Sewered population Urban land Other land Cultivated land	Bcol yr -1 93.5 person-1 425.1 ha-1 0.69 ha-1 60.4 ha-1	Sources: Sewered pop. Urban land Other land Confined waste Unconfined waste	Bcol yr -1 89.2 person -1 344.0 ha-1 0.20 ha-1 3.25 kgN-1 1.77 kgN-1
///////////////////////////////////////	o <u>ort:</u> (- coef.) (+ coef.)	Land-to-Water Tran Soil permeability Drainage density	(- coef.)
In-stream loss (day <sup>-1</sup> ) 0.13, 0.53, 0.64		In-stream loss (day 0.13, 0.49, 0.6	<del></del> ///////////////////////////////////
Reservoir loss (m yr1	70.6	Reservoir loss (m yr	<del>-1)</del> 79.0
R-squared	0.79	R-squared	0.80

Reach-accuracy

+/- 139%



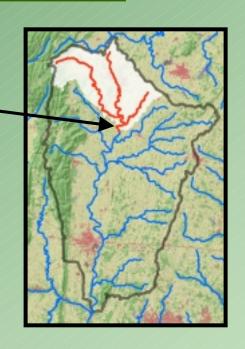
Reach-accuracy

#### **SPARROW SAS Predictions**

1. Flux at end of reach from total drainage —

PLOAD\_"SOURCES"

Units=mass per time



	🙀 predi	predict_load_lu3c5_update10.sas7bdat							
		WATERID	PLOAD_TOTAL	PLOAD_SEWERPOP	PLOAD_RESLAND	PLOAD_CONF	PLOAD_UNCONF	PLOAD_URBAN	
	1	1	121484650.0	66219466.197	12125206.853	5168037.222	3443474.063	34528465.718	
	2	2	120829611.6 67246411.49		12328268.739 4890852.6593	3240738.323	33123340.393		
	3	3	104455083.4	58110091.823	11945150.238	4126711.4436	2697056.243	27576073.743	
1	4	4	34836662.65	11209250.91	12100062.211	955631.90257	410579.0838	10161138.549	
	5	5	33461.09717	8815.981905	14806.559819	4470.8141532	3177.512838	2190.2284615	



#### **SPARROW SAS Predictions**

2. Flux delivered to end of reach w/o aquatic decay.

PLOAD\_ND\_TOTAL PLOAD\_ND\_"SOURCES"

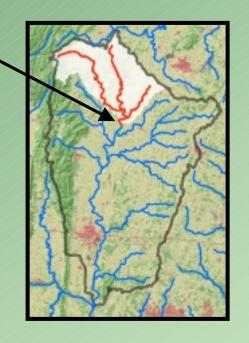
Units=mass per time

Total mass removed in streams & reservoirs =

PLOAD\_ND\_TOTAL - PLOAD\_TOTAL

mass removed in stream = (PLOAD\_ND\_TOTAL

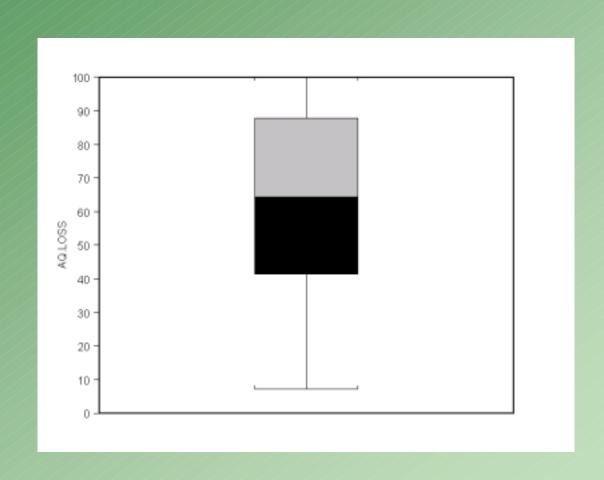
- PLOAD\_TOTAL) / PLOAD\_ND\_TOTAL \* 100



pred	predict_load_lu3c5_update10.sas7bdat									
	PLOAD_ND_TOTAL	PLOAD_ND_SEWERPOP	PLOAD_ND_RESLAND	PLOAD_ND_CONF	PLOAD_ND_UNCONF	PLOAD_ND_URBAN				
1	212308549.82	96302794.235	28477245.55	13469327.827	8812938.0104	65246244.192				
2	207354644.69	95473289.756	28342708.098	12922726.18	8424455.3553	62191465.303				
8	190271689.97	85893850.93	27907222.722	12128509.75	7859987.0468	56482119.517				
4	53201191.121	14699176.151	21403030.922	2054432.0414	700102.92756	14344449.079				
5	33853.632403	8919.4030041	14980.256942	4523.2616876	3214.7885356	2215.9222342				



# SPARROW Fecal Coliform Model Aquatic Loss (305 NASQAN sites)





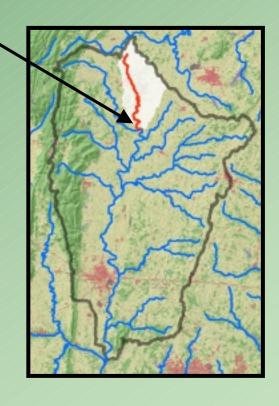
#### **SPARROW SAS Predictions**

3. Flux delivered to reach from incremental reach catchment

INC\_TOTAL INC\_"SOURCES"

Units = mass per time

	predict_load_lu3c5_update10.sas7bdat							
		INC_TOTAL	INC_SEWERPOP	INC_RESLAND	INC_CONF	INC_UNCONF	INC_URBAN	
ı	1	117402.79	53516.418025	1727.7574909	9987.6667	7098.469835	45072.478	
ı	2	2690867.4	847343.28539	3413.5396544	18282.879	12994.07299	1808833.6	
ı	3	1915108.9	196226.86609	446.97982087	0	0	1718435.1	
ı	4	6927415.4	2675820.9012	158071.72354	182251.45	129530.3984	3781740.9	
ı	5	33853.632	8919.4030041	14980.256942	4523.2616	3214.788535	2215.9222	
Ш					<del></del>			





### **SPARROW SAS Software**

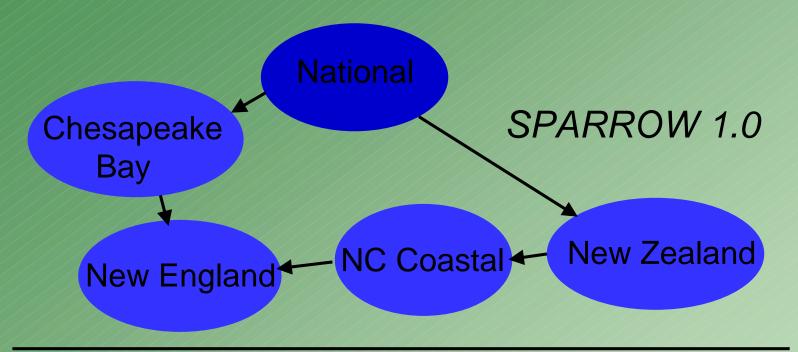
Execution run times for new SPARROW code:

New Zealand Waikato (5,000 reaches) ~ 0.30 minutes

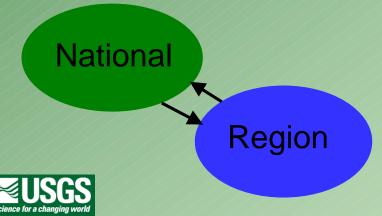
National (65,000 reaches) ~ 10 minutes



#### **Evolution of SPARROW Calibration Software**



#### SPARROW 2.0



#### SAS calibration / prediction software

- New revisions completed and on-going to national code
- Document and support single source of software maintained by the national SPARROW group

GIS methods and software –variety of approaches—will likely continue

#### **SPARROW SAS Software Enhancements**

#### Remain to be tested (\*\* = needs to be implemented)

- 1. Delivered flux (incremental, total, sources)
- 2. Parameter and prediction bootstrapping
- 3. Prediction confidence intervals
- 4. Model diagnostics:
  - a. Leverage statistic
  - b. Standardized residuals
  - c. Variance inflation factors (VIFs) \*\*
  - d. Parameter correlation matrix \*\*
  - e. Influence statistics (Cooks D) \*\*
  - f. Mallow's Cp \*\*
- 5. Weighted observations
  - -- "measurement error" -- load estimation error
  - --Hetereoscedasticity \*\*

